3D1 heavy chain variable region sequence

د می د	ATG M	GGT G	TGG W	AAC N	TGT C	ATC I	ATC I	TTC F	TTT F	30 CTG L	GTT V	ACA T	ACA T	GCT A	ACA T	GGT G	GTG V	CAC H	TCC S	60 CAG <u>Q</u>
	GTC V	CAG Q	CTG L	CAG Q	CAG Q	TCT S	GGG G	CCT P	GAG E	90 CTG L	GTG V	AGG R	CCT P	GGG G	GAA E	TCA S	GTG V	AAG K	ATT I	120 TCC S
	TGC C	AAG K	GGT G	TCC S	GGC G	TAC Y	ACA T	TTC F	ACT T	150 GAT D	TAT	GCT A	ATA I	CAG O	' TGG W	GTG V	AAG K	CAG Q	AGT S	180 CAT H
-	GCA A		AGT S		GAG E	TGG W	ATT I	GGA G	GTT V	210 ATT I	AAT N	ATT I	TAC Y	TAT Y	GAT D	AAT N	ACA T	AAC N	TAC Y	240 AAC <u>N</u>
<u> </u>	CAG O	AAG K	TTT F	AAG K	GGC G	AAG K	GCC A	ACA T	ATG M	270 ACT T	GTA V	GAC D	AAA K	TCC S	TCC S	AGC S	ACA T	GCC A	TAT Y	300 ATG M
	GAA E	CTT L	GCC A	AGA R	TTG L	ACA T	TCT S	GAG E	GAT D	330 TCT S	GCC A	ATC I	TAT Y	TAC Y	TGT C	GCA A	AGA R	GCG _A	GCC A	360 TGG W
	TAT Y	ATG M	GAC D	TAC Y	TGG W	GGT G	CAA Q	GGA G	ACC T	390 TCA S	GTC V	ACC T	GTC V	TCC S	TCA S		•			
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3D1 light chain variable region sequence

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GAC	ATT	GTG	CTG	TCA	CAG	TCT	CCA	TCC			GCT	GTG	TCA	GCA	GGA	GAG	AAG	GTC	ACT
₽	I	V	L	S	Q	s	P	s	s	L	A	V	S	A	G	E	K	v	T
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"W	Y.	Q	Q	K	P	G	Q	S	P	K	L	L	I	Y	W	A	S	Т	R
GAA	TCT	GGG	GTC	ССТ	GAT	C CC	mm c		270										300
E E	S	G	V	P	D	CGC R	TTC F	ACA T	GGC G	AGT	GGA	TCT	GGG		GAT	TTC	ACT	CTC	ACC
		Ü	•	-	ט	Λ	F	I.	G	S	G	·S	G	T	D	F	${f T}$	L	T
									330										360
ATC	AGC	AGT	GTG	CAG	GCT	GAA	GAC	CTG	GCA	GTT	TAT	TAC	TGC	ACG	CAA	TCT	TAT	AAT	CTT
I	S	S	V	Q	Α	E	D	L	A	V	Y	Y	C	T	0	S	Y	N	<u>L</u>
TAC	ACG	ጥጥር	GGA	GGG	GGG	200	3 3 C	0.00	390										
Y	ACG T	F	GGA	G	G	ACC . T	AAG K	CTG L	GAA E	ATA I	AAA								
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Figure 1 (B)

Hu3D1 heavy chain variable region sequence

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M	G	W	N	С	I	I	F	F	L	V	${f T}$	T	A	. Т	G	V	H	s	Ω
																			150
									90										120
GTC	CAG	CTG	GTG	CAG	TCT	GGG	GCT	GAG	GTG	AAG	AAG	CCI	' GGG	AGC	TCA	GTG	AAG	GTG	TCC
V	Q	L	V	Q	S	G	А	E	V	K	K	P	G	S	· S	V	K	V	s
									150										180
	AAA		TCC			ACA						ATA				AGA	CAG	GCT	CCT
С	K	A	S	G	Y	T	F	${f T}$	<u>D</u>	Y_	A_	<u> </u>	0	W	V	R	Q	A	P
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⊒ =CAG	AAG	TTT	AAG	GGC	AAG	GCC	ACA	ATG		СТА	GAC	מממ	ጥርር	ACG	AGC	ACA	GCC	TAT	ATG
<u> </u>	К	F	K	G	K	A	T	М	T	V	D	K	s	T	S	T	A	Y	M
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난 문									330										360
GAA	CTT	AGT	TCT	TTG	AGA	TCT	GAG	GAT	ACG	GCC	GTT	ŤАТ	TAC	TGT	GCA	AGA	GCG	GCC	TGG
ΨE	L	s	S	L	R	S	E	D	${f T}$	A	v	Y	Y	С	A	R	A	A	W
년 스																			
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	ATG	GAC	TAC	TGG	GGT	CAA	GGT	ACC	CTT	GTC	ACC	GTC	TCC	TCA					
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Hu3D1 light chain variable region sequence

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										150								•	•	
	TTA	AGC	TGC	AAA	TCC	AGT	CAG	AGT	CTG	CTC	220	ልርጥ	ACA	N.C.C	CCS	03.0				180
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	•										 -				R	E_	N_	<u>Y</u>	L	<u>A</u>
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-	rgg	TAC	CAG	CAG	AAA	CCA	GGG	CAG	CCT	CCA	מממ	CTG	CTC	አ ሙር	m» a	maa				240
_	W	Y	Q	. Q	K	P	G	Q	P	. p	K	L	L	I						
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Competition Binding Assay of Anti-B7.2 mAbs

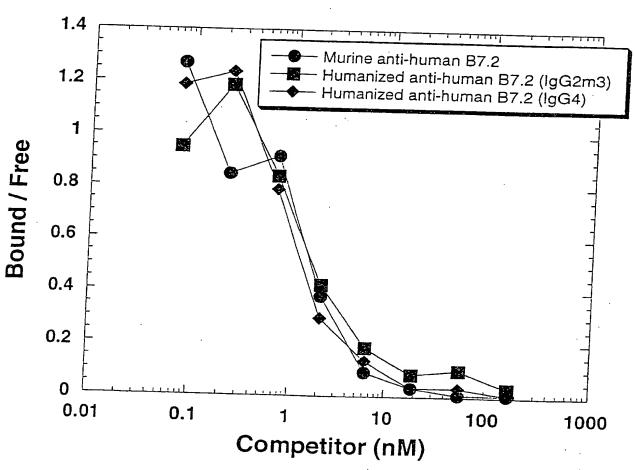


Figure 3

Direct Binding Assay of Anti-B7.2 mAbs

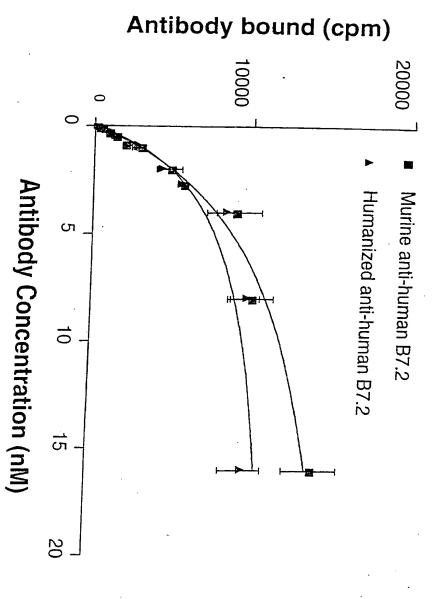


Figure 4

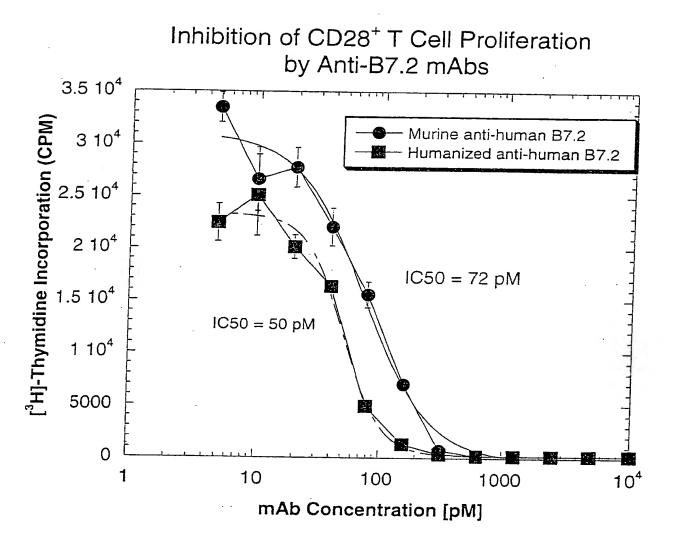
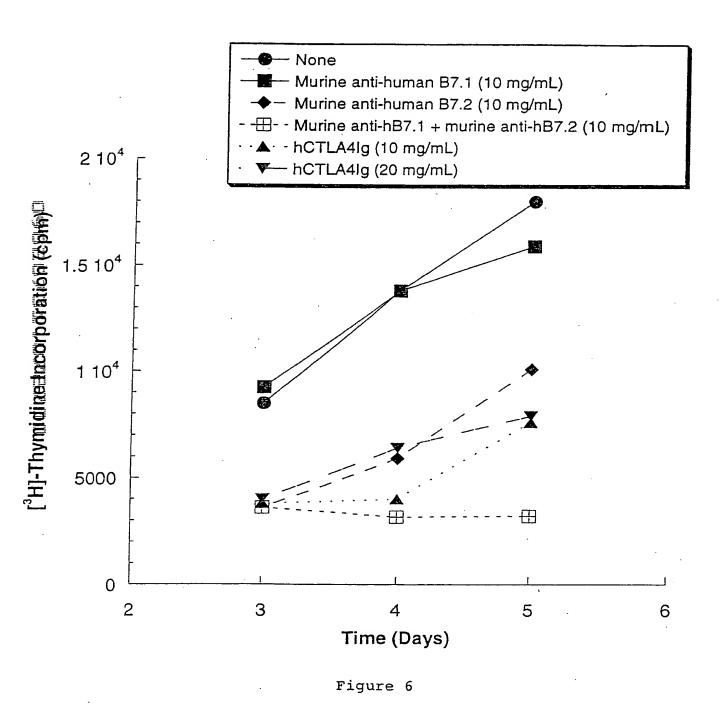


Figure 5

Inhibition of a Mixed Lymphocyte Reaction by Anti-B7 Antibodies and CTLA4IG



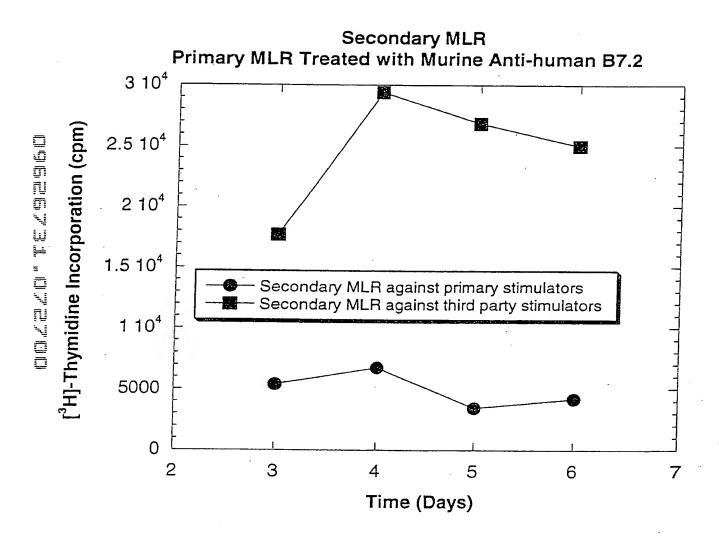


Figure 7

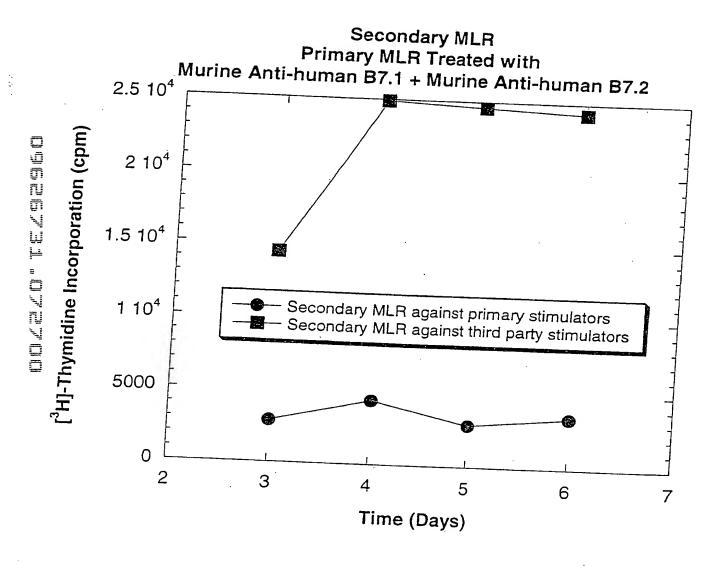
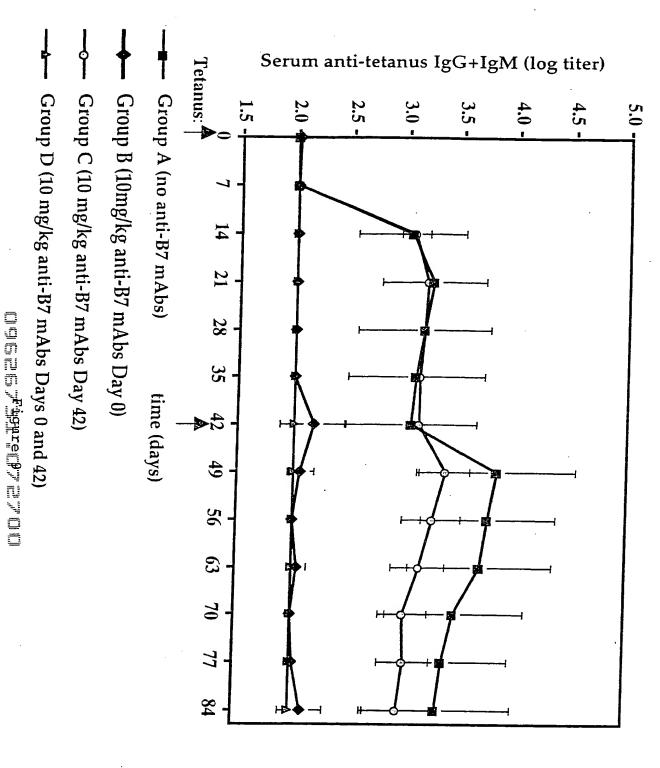


Figure 8

Antibody Response to Tetanus Immunization During Costimulation Blockade with Humanized Anti-B7.1 and Anti-B7.2



Anti-B7-2, Serum concentration

Serum Concentration of Humanized Anti-B7-2 in cynomolgus Monkeys

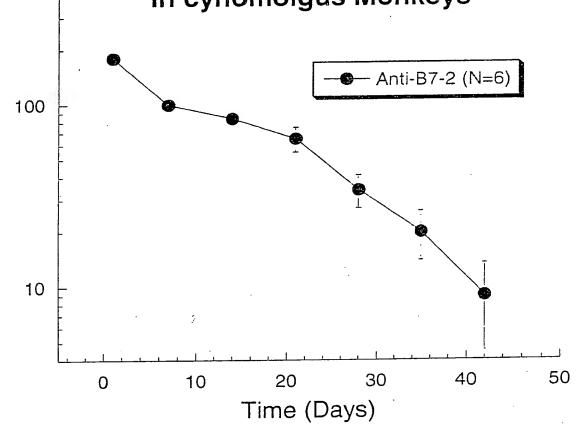


Figure 10